

d hist

(FILE 'HOME' ENTERED AT 12:07:57 ON 13 SEP 2006)

FILE 'REGISTRY' ENTERED AT 12:08:43 ON 13 SEP 2006

L1           STRUCTURE UPLOADED  
L2           18 S L1  
L3           448 S L1 FULL  
L4           0 S L3 AND MXS/CI

FILE 'CAPLUS' ENTERED AT 12:10:54 ON 13 SEP 2006

L5           214 S L3  
L6           30 S L5 AND (DIASTEREOM? OR ENANTIOMER?)  
L7           6 S L6 AND MIX?  
L8           16 S L6 AND PY<1997  
L9           5 S L7 AND PY<1997

FILE 'STNGUIDE' ENTERED AT 12:12:20 ON 13 SEP 2006

FILE 'CAPLUS' ENTERED AT 13:09:08 ON 13 SEP 2006

FILE 'STNGUIDE' ENTERED AT 13:09:10 ON 13 SEP 2006

FILE 'CAPLUS' ENTERED AT 13:16:09 ON 13 SEP 2006  
L10          STRUCTURE UPLOADED  
              S L10

FILE 'REGISTRY' ENTERED AT 13:29:33 ON 13 SEP 2006  
L11          50 S L10

FILE 'CAPLUS' ENTERED AT 13:29:34 ON 13 SEP 2006  
L12          44 S L11

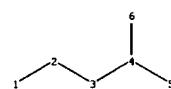
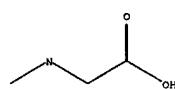
FILE 'REGISTRY' ENTERED AT 13:29:46 ON 13 SEP 2006  
L13          SCREEN 2051  
L14          SCREEN 2127  
L15          QUE L10 AND (L13 OR L14)  
L16          50 S L15  
L17          QUE L10 AND L13  
L18          50 S L17  
L19          SCREEN 1913  
L20          QUE L10 NOT L19  
L21          50 S L20  
L22          162357 S L20 FULL

FILE 'CAPLUS' ENTERED AT 13:50:38 ON 13 SEP 2006  
L23          314332 S L22  
L24          2544 S L23 AND (COMBINATOR? OR LIBRAR?)  
L25          54 S L24 AND (ENANTIOMER? OR DIASTEREOM?)  
L26          4 S L25 AND PY<1997

FILE 'REGISTRY' ENTERED AT 13:54:56 ON 13 SEP 2006  
L27          10 S 136083-74-4 OR 35899-43-5 OR 181213-21-8 OR 181213-23-0 OR 18

FILE 'REGISTRY' ENTERED AT 14:15:35 ON 13 SEP 2006  
L28          10 S 136083-74-4 OR 35899-43-5 OR 181213-21-8 OR 181213-23-0 OR 18

FILE 'CAPLUS' ENTERED AT 14:16:41 ON 13 SEP 2006  
L29          367 S L24 AND PY<1997  
L30          0 S L29 AND (MANNICH? OR REDUCTIV(2N)AMINAT? OR IMMIN? OR ALDIMIN  
L31          247 S L24 AND PY<1996



chain nodes :

1 2 3 4 5 6

chain bonds :

1-2 2-3 3-4 4-5 4-6

exact/norm bonds :

1-2 2-3

exact bonds :

3-4

normalized bonds :

4-5 4-6

Connectivity :

3:1 M minimum RC ring/chain

Match level :

1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS